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10/660,434	09/10/2003	Guennadi V. Glinskii	23543-07570	4883
758 FENWICK & V	7590 08/07/200 VEST LLP	EXAMINER		
SILICON VAL	LEY CENTER	LIN, JERRY		
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Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

	Application No.	Applicant(s)			
	10/660,434	GLINSKII, GUENNADI V.			
Office Action Summary	Examiner	Art Unit			
	JERRY LIN	1631			
The MAILING DATE of this communication app Period for Reply	ears on the cover sheet with the c	orrespondence address			
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period w. - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tim vill apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	lely filed the mailing date of this communication. (35 U.S.C. § 133).			
Status					
1) Responsive to communication(s) filed on <u>05 Ma</u>	action is non-final. nce except for formal matters, pro				
Disposition of Claims					
4) ☐ Claim(s) 1-28 and 36-38 is/are pending in the a 4a) Of the above claim(s) is/are withdrav 5) ☐ Claim(s) is/are allowed. 6) ☐ Claim(s) 1-6,8-13 and 16-228 is/are rejected. 7) ☐ Claim(s) 7,14 and 15 is/are objected to. 8) ☐ Claim(s) are subject to restriction and/or Application Papers 9) ☐ The specification is objected to by the Examinet 10) ☐ The drawing(s) filed on is/are: a) ☐ access Applicant may not request that any objection to the orection and or declaration is objected to by the Examinet Replacement drawing sheet(s) including the correction in the orection of the orection and or declaration is objected to by the Examinet and or declaration is objected to be an or declaration is objected to be an or declaration in the order and or declaration is objected to be an order and or declaration is objected to be an order and order and order and order and or	r election requirement. r. epted or b) □ objected to by the Edrawing(s) be held in abeyance. See ion is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan in the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required if the drawing(s) is objected to by the legan is required in th	e 37 CFR 1.85(a). ected to. See 37 CFR 1.121(d).			
	animor. Noto the attached office	7.00.001 01 101111 1 0 102.			
Priority under 35 U.S.C. § 119 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some color None of: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No. 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received.					
Attachment(s) 1) Notice of References Cited (PTO-892) 2) Notice of Draftsperson's Patent Drawing Review (PTO-948) 3) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date 7/14/08.	4) Interview Summary Paper No(s)/Mail Da 5) Notice of Informal P 6) Other:	ite			

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DETAILED ACTION

1. Applicants' arguments, filed May 5, 2008, have been fully considered and they are deemed to be persuasive. The following rejections are newly applied. They constitute the complete set presently being applied to the instant application.

Status of the Claims

Claims 1-28 and 36-38 are under examination.

Regarding claims 24-28, the Table 69 was elected as species, and the other tables are withdrawn as being drawn to a non-elected species.

Claim Rejections - 35 USC § 103

- 2. The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:
 - (a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.
- 3. Claims 1-6, 8-11, and 16-23 are rejected under 35 U.S.C. 103(a) as being unpatentable over Alizadeh et al. (Nature (2000) Volume 403, pages 503-511) in view of Bertucci et al. (Human Molecular Genetics (2000) Volume 9, Number 20, pages 2981-2991).

The instant claims are drawn to identifying a subset of genes using a correlation coefficient calculated from gene expression data.

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Regarding claim 1, Alizadeh et al. teach using two samples that different with respect to phenotype and determining a reference set of genes (page 504, section under "Analysis of gene expression in lymphoid malignancies"), identifying a second reference set of expressed genes independent of the first reference set from a third and fourth sample where the third or fourth sample is from a different source than the first and second sample (page 504, section under "Analysis of gene expression in lymphoid malignancies"); identifying a concordance set of expressed genes with the same direction of differential expression (i.e. identifying genes common to the first and second reference sets) (page 504 - page 505, section under "Gene expression patterns and tumor phenotype"). Furthermore, Alizadeh et al. recorded the subset of genes within a concordance set in a user-readable format (page 504, section under "Gene expression patterns and tumor phenotype" and Figure 2).

However, Alizadeh et al. do not teach determining a correlation coefficient that exceeds a predetermined value.

Regarding claim 1, Bertucci et al. teaches determining the correlation coefficient that exceed a predetermine value for correlating genes (page 2987, right column, 2nd full paragraph).

Regarding claims 2-6, Bertucci et al. teach determining a correlation coefficient (page 2987, right column, 2nd full paragraph); logarithmically transforming the differentials (page 2987, right column, 3rd full paragraph); wherein the correlation

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coefficient has an absolute value greater than 0.98 (page 2987, right column, 2nd full paragraph).

Regarding claims 8-11, Bertucci et al. teach wherein the gene expression data is cDNA or RNA quantification data (page 870); wherein the sample comprises of a cell line, which is a tumor cell line (page 869, left column, top).

Regarding claims 16-19, Bertucci et al. teach wherein the sample is from a patient, a healthy donor, is a tumor cell, or from the colon (paragraph bridging pages 869-867).

Regarding claim 20, Bertucci et al. teach where the phenotype is selected from lymph node status (page 2983, left column, bottom section).

Regarding claim 21, Bertucci et al. teach where a plurality of independent samples is used for each sample (paragraph bridging page 869-870); and where the differential is an average over the sample (page 870, Table II).

Regarding claim 22, Bertucci et al. teach determining a second correlation coefficient with a positive sign that establishes a positive correlation with a phenotype (page 2983, right column, first full paragraph).

Regarding claim 23, Bertucci et al. teach that ERBB2 had the highest correlation of the genes tested (page 2983, first full paragraph) and since its correlation was the highest its correlation was the most reliable indication of cancer (abstract; page 2985, left column, bottom paragraph – right column, top paragraph).

It would have been obvious to one of ordinary skill in the art at the time of the invention to combine the methods of Alizadeh et al. with Bertucci et al. to gain the

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advantage of determining the reproducibility of experiments. One of the challenges that face gene expression experiments is the precision of the equipment used in those experiments. Given the same experiment, the equipment may produce different results. Given the uncertainty of the equipment as well as the need to compare data from different sources, one of ordinary skill in the art would seek to verify the reproducibility of experiments to ensure that their interpretation of the data is correct. Bertucci et al. teaches a method of determining the accuracy of gene expression using a correlation coefficient (page 2987, right column, 2nd full paragraph). Thus, one of ordinary skill in the art would be motivated to combine the methods of Alizadeh et al. and Bertucci et al. to ensure that the gene expression data was accurate.

Response to Arguments

4. Although this is a new rejection, Applicants have responded the citation of Bertucci et al. in a previous rejection by stating that Bertucci et al.'s methods are used to verify experimental results and measure reproducibility of differential expression.

Applicants also state that is unclear how a method used for verifying experimental results and measuring reproducibility of differential expression may be used to select a subset of genes. Although, Bertucci et al. verifies experimental results and measure reproducibility of differential expression, their method also determines "Correlations of gene pairs based on expression profiles were measured with the correlation coefficient r." (page 2987, right column, bottom paragraph). Thus, once the subset of genes is identified as a concordance set, Bertucci et al.'s method applies to determine the subset

of genes correlated between different samples or reference sets (page 2987, right column, bottom paragraph).

5. Claims 12 and 13 are rejected under 35 U.S.C. 103(a) as being unpatentable over Alizadeh et al. (Nature (2000) Volume 403, pages 503-511) in view of Bertucci et al. (Human Molecular Genetics (2000) Volume 9, Number 20, pages 2981-2991) further in view of Young et al. (US #2005/0255588 A1).

The instant claims are drawn to identifying a subset of genes using a correlation coefficient calculated from gene expression data and using samples comprising omnipotent or pluripotent cells.

Alizadeh et al. and Bertucci et al. are applied as above.

However, neither Alizadeh et al. nor Bertucci et al. teach using pluripotent or omnipotent stem cells.

Young et al. teach creating samples of pluripotent or omnipotent stem cells (page 3, paragraph 0019-0020).

It would have been obvious to one of ordinary skill in the art at the time of the invention to use the cell lines described in Young et al. with the methods of Alizadeh et al. and Bertucci et al. The motivation to combine Alizadeh et al. with Bertucci et al. is provided above. It is recognized in the art the use of stem cells could potentially provide many new applications in science and medicine (Young et al., page 3, paragraph 0015-0016). However, it is unclear what genes are expressed to maintain a cell as a stem cell. Thus one of ordinary skill in the art seeking to understand the gene expression of

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stem cells would be motivated to determine what genes are differentially expressed that correspond to the stem cell phenotype. Alizadeh et al. provide a method of identifying the correct identification of the differences in gene expression between different cell lines (Alizadeh et al., page 504). Alizadeh et al.'s method could identify the differences in gene expression in a stem cell as compared to other cells. Thus one of ordinary skill in the art seeking to understand the gene expression of stem cells would be motivated to take the sample cell lines disclosed by Young et al. and use those samples in the method provided by Alizadeh et al.

6. Claims 24-28 and 36-38 are rejected under 35 U.S.C. 103(a) as being unpatentable over Alizadeh et al. (Nature (2000) Volume 403, pages 503-511) in view of Bertucci et al. (Human Molecular Genetics (2000) Volume 9, Number 20, pages 2981-2991) in view of Horne et al. (US 6,974,667) in view of Dai et al. (US 7,171311) in view of Ohara et al. (DNA Res. (1997) Volume 4, number 5, pages 345-349) in view of Bach (Nucleic Acids Research (1991) Volume 19, number 13, pages 3553-3559) in view of Volta et al. (Genomics (1999) Volume 55, number 1, pages 68-77) in view of Ohara et al. (DNA Res. (1998) Volume 5, number 3, pages 169-176) in view of Venter et al. (US 6,812,339), in view of Calabretta et al. (US 5,734,039), in view of Wohlgemuth et al. (US 6,905,827), in view of Cocks et al. (US 6,607,879).

Alizadeh et al. and Bertucci et al. are applied as above.

However, neither Barney et al. nor Blasband et al. teach using the genes listed in Table 69.

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Horne et al. disclose the genes with the sequences of Seq ID 2240 (MGC5466), Seq ID 2249 (MGC5466), Seq ID 2254 (IER3) (see sequence search report).

Dai et al. disclose the genes with the sequences of Seq ID 2241 (Wnt5A), Seq ID 2245 (ITPR1), Seq ID 2250 (CHAF1A) (see sequence search report).

Ohara et al. (1997) discloses the gene AB007945 (i.e., KIAA0476, Seq ID 2243) (see NCBI printout).

Bach discloses the gene with the sequence of Seq ID 2247 (TCF2) (see sequence search report).

Volta et al. discloses the gene with the sequence of Seq ID 2252 (CDS2) (see sequence search report).

Ohara et al. (1998) discloses the gene of Seq ID 2256 (PPFIA3) (see ncbi search results).

Venter et al. discloses the gene of Seq ID 2258 (COPEB) (see sequence search report).

Calabretta et al. discloses the gene of Seq ID 2260 (FOS) (see sequence search report).

Wohlgemuth et al. discloses the gene of Seq ID 2262 (JUNB) (see sequence search report).

Cocks et al. discloses the gene of Seq ID 2264 (ZFP36) (see sequence search report).

It would have been obvious to one of ordinary skill in the art to combine the methods of Alizadeh et al. and Bertucci et al. with the probes for the genes of Horne et

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al. Dai et al., Ohara et al., Venter et al., Wohlgemuth et al. and Cocks et al. Each of the genes of Table 69 are known in the prior art as shown above. Given that Alizadeh et al. and Bertucci et al. both analyze gene expression levels on microarrays, one of ordinary skill in the art at the time of the invention could have substituted the genes used by Alizadeh et al. and Bertucci et al. with the genes of Table 69 for the predictable result of analyzing the gene expression levels of the genes of Table 69.

Allowable Subject Matter

7. Claims 7, 14 and 15 are objected to as being dependent upon a rejected base claim, but would be allowable if rewritten in independent form including all of the limitations of the base claim and any intervening claims.

Withdrawn Rejections

8. Applicant's arguments and amendments, filed 5/05/08, with respect to the rejection(s) of claim(s) 24-28 and 36-38 under 35 U.S.C. §112 1st and 2nd Paragraph have been fully considered and are persuasive. Applicants have demonstrated how the sequence ID number correspond to the genes in Table 69 and amended the specification to clarify the correspondence. Applicants have also shown GenBank IDs listed in Table 69 can be found in the "Gene" database on the NCBI website. Therefore, these rejections have been withdrawn.

Applicant's arguments and amendments, filed 5/05/08, with respect to the rejection(s) of claim(s) 1-28 and 36-38 under 35 U.S.C. §101 Paragraph have been fully

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considered and are persuasive. The amendment to the claim is sufficient to overcome this rejection. This rejection has been withdrawn.

Applicant's arguments and amendments, filed 5/05/08, with respect to the rejection(s) of claim(s) 1-23 under 35 U.S.C. §103 as being unpatentable over Backert et al. in view of Bertucci et al. have been fully considered and are persuasive. Neither Backert et al. nor Bertucci et al. teach the new limitation of a second reference set of expressed genes that includes at least one gene that is not included in the first reference set. The amendment to the claim 1 is sufficient to overcome this rejection. This rejection has been withdrawn.

Contact Information

Any inquiry concerning this communication or earlier communications from the examiner should be directed to JERRY LIN whose telephone number is (571)272-2561. The examiner can normally be reached on 7:00-5:30pm, M-TH.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Marjorie A. Moran can be reached on (571) 272-0720. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

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Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

/Jerry Lin/ Examiner, Art Unit 1631 7/31/08